

## **RAW SEQUENCE LISTING**

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Application Serial Number: 09/674,8170  
Source: IFW/6  
Date Processed by STIC: 4/22/05

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IFW16

## RAW SEQUENCE LISTING

DATE: 04/22/2005

PATENT APPLICATION: US/09/674,817D

TIME: 12:10:44

Input Set : A:\51441349.app

Output Set: N:\CRF4\04222005\I674817D.raw

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4 <110> APPLICANT: LORZ, HORST
5      LUTTICKE, STEPHANIE
6      ABEL, GERNOT
7      GENSCHER, ULRICH
9 <120> TITLE OF INVENTION: NUCLEIC ACID MOLECULES WHICH CODE FOR ENZYMES
10     DERIVED FROM WHEAT AND WHICH ARE INVOLVED IN
11     THE SYNTHESIS OF STARCH
13 <130> FILE REFERENCE: 514413-3849.1
15 <140> CURRENT APPLICATION NUMBER: 09/674,817D
16 <141> CURRENT FILING DATE: 2000-11-06
18 <150> PRIOR APPLICATION NUMBER: WO 99/58690
19 <151> PRIOR FILING DATE: 1999-11-18
21 <150> PRIOR APPLICATION NUMBER: DE 198 20 608.9
22 <151> PRIOR FILING DATE: 1998-05-08
24 <160> NUMBER OF SEQ ID NOS: 11
26 <170> SOFTWARE: PatentIn version 3.3
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 2997
30 <212> TYPE: DNA
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48   1              5              10              15
50 gcg ggg aag ggg gtc gcc gag gtg tgc gcc gcg gtt gtc gag gcg gcg      95
51 Ala Gly Lys Gly Val Gly Glu Val Cys Ala Ala Val Val Glu Ala Ala
52              20              25              30
54 acg aag gta gag gac gag ggg gag gag gac gag ccg gtg gcg gag gac      143
55 Thr Lys Val Glu Asp Glu Gly Glu Glu Asp Glu Pro Val Ala Glu Asp
56              35              40              45
58 agg tac gcg ctc gcc gcc gcg tgc agg gtg ctc gcc gga atg ccc gcg      191
59 Arg Tyr Ala Leu Gly Gly Ala Cys Arg Val Leu Ala Gly Met Pro Ala
60              50              55              60
62 ccg ctg gcc gcc acc gcg ctc gcc gcc ggg gtc aat ttc gcc gtc tat      239

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66 tcc ggc gga gcc acc gcc gcg gcg ctc tgc ctc ttc acg cca gaa gat      287
67 Ser Gly Gly Ala Thr Ala Ala Ala Leu Cys Leu Phe Thr Pro Glu Asp
68 80              85              90              95
70 ctc aag gcg gtgggggttc ctcccgagta gagttcatca gctttgcgtg      336
71 Leu Lys Ala
74 cgccgcgcgc cctttttttg ggccctgcaat ttaagttttg tactgggggca aatgctgcag      396
76 gat agg gtg acc gag gag gtt ccc ctt gac ccc ctg atg aat cgg acc      444
77 Asp Arg Val Thr Glu Glu Val Pro Leu Asp Pro Leu Met Asn Arg Thr
78      100              105              110
80 ggg aac gtg tgg cat gtc ttc atc gaa ggc gag ctg cac aac atg ctt      492
81 Gly Asn Val Trp His Val Phe Ile Glu Gly Glu Leu His Asn Met Leu
82 115              120              125              130
84 tac ggg tac agg ttc gac ggc acc ttt gct cct cac tgc ggg cac tac      540
85 Tyr Gly Tyr Arg Phe Asp Gly Thr Phe Ala Pro His Cys Gly His Tyr
86              135              140              145
88 ctt gat gtt tcc aat gtc gtg gtg gat cct tat gct aag gca gtg ata      588
89 Leu Asp Val Ser Asn Val Val Val Asp Pro Tyr Ala Lys Ala Val Ile
90              150              155              160
92 agc cga ggg gag tat ggt gtt cca gcg cgt ggt aac aat tgc tgg cct      636
93 Ser Arg Gly Glu Tyr Gly Val Pro Ala Arg Gly Asn Asn Cys Trp Pro
94      165              170              175
96 cag atg gct ggc atg atc cct ctt cca tat agc acg ttt gat tgg gaa      684
97 Gln Met Ala Gly Met Ile Pro Leu Pro Tyr Ser Thr Phe Asp Trp Glu
98      180              185              190
100 ggc gac cta cct cta aga tat cct caa aag gac ctg gta ata tat gag      732
101 Gly Asp Leu Pro Leu Arg Tyr Pro Gln Lys Asp Leu Val Ile Tyr Glu
102 195              200              205              210
104 atg cac ttg cgt gga ttc acg aag cat gat tca agc aat gta gaa cat      780
105 Met His Leu Arg Gly Phe Thr Lys His Asp Ser Ser Asn Val Glu His
106              215              220              225
108 ccg ggt act ttc att gga gct gtg tcg aag ctt gac tat ttg aag gag      828
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112 ctt gga gtt aat tgt att gaa tta atg ccc tgc cat gag ttc aac gag      876
113 Leu Gly Val Asn Cys Ile Glu Leu Met Pro Cys His Glu Phe Asn Glu
114              245              250              255
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117 Leu Glu Tyr Ser Thr Ser Ser Ser Lys Met Asn Phe Trp Gly Tyr Ser
118      260              265              270
120 acc ata aac ttc ttt tca cca atg aca aga tac aca tca ggc ggg ata      972
121 Thr Ile Asn Phe Phe Ser Pro Met Thr Arg Tyr Thr Ser Gly Gly Ile
122 275              280              285              290
124 aaa aac tgt ggg cgt gat gcc ata aat gag ttc aaa act ttt gta aga      1020
125 Lys Asn Cys Gly Arg Asp Ala Ile Asn Glu Phe Lys Thr Phe Val Arg
126              295              300              305
128 gag gct cac aaa cgg gga att gag gtg atc ctg gat gtt gtc ttc aac      1068
129 Glu Ala His Lys Arg Gly Ile Glu Val Ile Leu Asp Val Val Phe Asn

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133 His Thr Ala Glu Gly Asn Glu Asn Gly Pro Ile Leu Ser Phe Lys Gly				
134	325	330	335	
136 gtc gat aat act aca tac tat atg ctt gca ccc aag gga gag ttt tat				1164
137 Val Asp Asn Thr Thr Tyr Tyr Met Leu Ala Pro Lys Gly Glu Phe Tyr				
138	340	345	350	
140 aac tat tct ggc tgt ggg aat acc ttc aac tgt aat cat cct gtg gtt				1212
141 Asn Tyr Ser Gly Cys Gly Asn Thr Phe Asn Cys Asn His Pro Val Val				
142 355	360	365	370	
144 cgt caa ttc att gta gat tgt tta aga tac tgg gtg acg gaa atg cat				1260
145 Arg Gln Phe Ile Val Asp Cys Leu Arg Tyr Trp Val Thr Glu Met His				
146	375	380	385	
148 gtt gat ggt ttt cgt ttt gat ctt gca tcc ata atg acc aga ggt tcc				1308
149 Val Asp Gly Phe Arg Phe Asp Leu Ala Ser Ile Met Thr Arg Gly Ser				
150	390	395	400	
152 agt ctg tgg gat cca gtt aac gtg tat gga gct cca ata gaa ggt gac				1356
153 Ser Leu Trp Asp Pro Val Asn Val Tyr Gly Ala Pro Ile Glu Gly Asp				
154	405	410	415	
156 atg atc aca aca ggg aca cct ctt gtt act cca cca ctt att gac atg				1404
157 Met Ile Thr Thr Gly Thr Pro Leu Val Thr Pro Pro Leu Ile Asp Met				
158	420	425	430	
160 atc agc aat gac cca att ctt gga ggc gtc aag ctc att gct gaa gca				1452
161 Ile Ser Asn Asp Pro Ile Leu Gly Gly Val Lys Leu Ile Ala Glu Ala				
162 435	440	445	450	
164 tgg gat gca gga ggc ctc tat caa gta ggt caa ttc cct cac tgg aat				1500
165 Trp Asp Ala Gly Gly Leu Tyr Gln Val Gly Gln Phe Pro His Trp Asn				
166	455	460	465	
168 gtt tgg tct gag tgg aat ggg aag tac cgg gac att gtg cgt caa ttc				1548
169 Val Trp Ser Glu Trp Asn Gly Lys Tyr Arg Asp Ile Val Arg Gln Phe				
170	470	475	480	
172 att aaa ggc act gat gga ttt gct ggt ggt ttt gcc gaa tgt ctt tgt				1596
173 Ile Lys Gly Thr Asp Gly Phe Ala Gly Gly Phe Ala Glu Cys Leu Cys				
174	485	490	495	
176 gga agt cca cac cta tac cag gtaagttgtg gcaatacttg taaatgagtt				1647
177 Gly Ser Pro His Leu Tyr Gln				
178	500	505		
180 gagtgaatgt cacctggatt ttttatatat accacatgat gatacacatc taaatatata				1707
182 acaatcatag tgtatgcata tgcatttggc taagaagtat tagtgtatac actagtgcata				1767
184 tatatagggt ttaacaccca acttgccaat gaaggaacat agggctttct agttatctta				1827
186 tttatttgtc cgggtgaataa tccactgaaa aattccagcc atgtcatttt ttagggggggg				1887
188 agaagaaact atattgattt gccccctaa aagaagccat ctcagaattc ataggtaagt				1947
190 tgcttttctg taaagaaagg aaaacgactt catactttct atcgggtgcta acttagctcg				2007
192 atgtatatatt gtaagatgaa tgccaaattt aatttgtcgg ataatttgat ctgttattca				2067
194 caaatttcta tttgttttct ctagaaatca aaccagtaac ttgttatttg cactgcaact				2127
196 tcttattgat taatcag gca gga gga agg aaa cct tgg cac agt atc aac				2177
197	Ala Gly Gly Arg Lys Pro Trp His Ser Ile Asn			
198	510	515		
200 ttt gta tgt gca cat gat gga ttt aca ctg gct gat ttg gta aca tat				2225

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205	Asn	Lys	Lys	Tyr	Asn	Leu	Pro	Asn	Gly	Glu	Asn	Asn	Arg	Asp	Gly	Glu	
206			535					540					545				
208	aat	cac	aat	ctt	agc	tgg	aat	tgt	ggg	gag	gaa	gga	gaa	ttc	gca	aga	2321
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210		550					555					560					
212	ttg	tct	gtc	aaa	aga	ttg	agg	aag	agg	cag	atg	cgc	aat	ttc	ttt	gtt	2369
213	Leu	Ser	Val	Lys	Arg	Leu	Arg	Lys	Arg	Gln	Met	Arg	Asn	Phe	Phe	Val	
214	565					570					575					580	
216	tgt	ctc	atg	gtt	tct	caa	gga	gtt	cca	atg	ttc	tac	atg	ggt	gat	gaa	2417
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220	tat	ggc	cac	aca	aaa	ggg	ggc	aac	aac	aat	aca	tac	tgc	cat	gat	tct	2465
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224	tat	gtc	aat	tat	ttt	cgc	tgg	gat	aaa	aaa	gaa	caa	tac	tct	gag	ttg	2513
225	Tyr	Val	Asn	Tyr	Phe	Arg	Trp	Asp	Lys	Lys	Glu	Gln	Tyr	Ser	Glu	Leu	
226			615					620					625				
228	cac	cga	ttc	tgc	tgc	ctc	atg	acc	aaa	ttc	cgc	aag	gag	tgc	gag	ggt	2561
229	His	Arg	Phe	Cys	Cys	Leu	Met	Thr	Lys	Phe	Arg	Lys	Glu	Cys	Glu	Gly	
230		630					635					640					
232	ctt	ggc	ctt	gag	gac	ttt	cca	acg	gcc	aaa	cgg	ctg	cag	tgg	cat	ggt	2609
233	Leu	Gly	Leu	Glu	Asp	Phe	Pro	Thr	Ala	Lys	Arg	Leu	Gln	Trp	His	Gly	
234	645					650				655						660	
236	cat	cag	cct	ggg	aag	cct	gat	tgg	tct	gag	aat	agc	cga	ttc	gtt	gcc	2657
237	His	Gln	Pro	Gly	Lys	Pro	Asp	Trp	Ser	Glu	Asn	Ser	Arg	Phe	Val	Ala	
238				665					670					675			
240	ttt	tcc	atg	aaa	gat	gaa	aga	cag	ggc	gag	atc	tat	gtg	gcc	ttc	aac	2705
241	Phe	Ser	Met	Lys	Asp	Glu	Arg	Gln	Gly	Glu	Ile	Tyr	Val	Ala	Phe	Asn	
242				680					685					690			
244	acc	agc	cac	tta	ccg	gcc	gtt	gtt	gag	ctc	cca	gag	cgc	gca	ggg	cgc	2753
245	Thr	Ser	His	Leu	Pro	Ala	Val	Val	Glu	Leu	Pro	Glu	Arg	Ala	Gly	Arg	
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249	Arg	Trp	Glu	Pro	Val	Val	Asp	Thr	Gly	Lys	Pro	Ala	Pro	Tyr	Asp	Phe	
250		710					715					720					
252	ctc	acc	gac	gac	tta	cct	gat	cgc	gct	ctc	acc	ata	cac	cag	ttc	tcg	2849
253	Leu	Thr	Asp	Asp	Leu	Pro	Asp	Arg	Ala	Leu	Thr	Ile	His	Gln	Phe	Ser	
254	725					730					735					740	
256	cat	ttc	ctc	tac	tcc	aac	ctc	tac	ccc	atg	ctc	agc	tac	tca	tcg	gtc	2897
257	His	Phe	Leu	Tyr	Ser	Asn	Leu	Tyr	Pro	Met	Leu	Ser	Tyr	Ser	Ser	Val	
258				745					750					755			
260	atc	cta	gta	ttg	cgc	cct	gat	gtt	tga	gag	acc	aat	ata	tac	agt	aaa	2945
261	Ile	Leu	Val	Leu	Arg	Pro	Asp	Val		Glu	Thr	Asn	Ile	Tyr	Ser	Lys	
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264	taa	tat	gtc	tat	atg	taaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	aaaaaaaaaaaa	2997
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